

Entrez Nucleotide

Nucleotide

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1: [AI817366](#) Reports Links
wk22d07.x1 NCI_CGAP_Lym12 Homo sapiens cDNA clone IMAGE:2413069 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|5436445|gb|AI817366.1|[5436445]

2: [AI797688](#) Reports Links
we90d08.x1 SoaresNFLT_GBC_S1 Homo sapiens cDNA clone IMAGE:2348367 3'
similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|5363160|gb|AI797688.1|[5363160]

3: [AI794615](#) Reports Links
fc46f02.y1 Zebrafish WashU MPIMG EST Danio rerio cDNA clone IMAGE:3724443 5'
similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|5342331|gb|AI794615.1|[5342331]

4: [AI762942](#) Reports Links
wh94f09.x1 NCI_CGAP CLL1 Homo sapiens cDNA clone IMAGE:2388425 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|5178609|gb|AI762942.1|[5178609]

5: [AI726943](#) Reports Links
BNLGH16897 Six-day Cotton fiber Gossypium hirsutum cDNA 5' similar to (AB012122)
TIP49 [Homo sapiens] gi|3243035, mRNA sequence
gi|5045795|gb|AI726943.1|[5045795]

6: [AI725799](#) Reports Links
BNLGH13195 Six-day Cotton fiber Gossypium hirsutum cDNA 5' similar to (AB012122)
TIP49 [Homo sapiens] gi|3243035, mRNA sequence
gi|5044651|gb|AI725799.1|[5044651]

7: [AI722028](#) Reports Links
fd18e04.y1 Sugano Kawakami zebrafish DRA Danio rerio cDNA clone IMAGE:2351934
5' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|5040357|gb|AI722028.1|[5040357]

8: [AI721333](#) Reports Links

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fd18e04.x1 Sugano Kawakami zebrafish DRA Danio rerio cDNA clone IMAGE:2351934
 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|5039662|gb|AI721333.1|[5039662]

9: AB002406 Reports

Rattus norvegicus mRNA for TIP49, complete cds
 gi|2225876|dbj|AB002406.1|[2225876]

Links

10: AI672922 Reports

we74b05.x1 Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone
 IMAGE:2346801 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|4852653|gb|AI672922.1|[4852653]

Links

11: AI659383 Reports

tul1f02.x1 NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2250747 3' similar to
 TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|4762953|gb|AI659383.1|[4762953]

Links

12: AB014075 Reports

Clostridium histolyticum genes for hypoxanthine-guanine phosphoribosyl-transferase
 (HGPRTase), GTPase and 12 ORFs, complete and partial cds
 gi|3868863|dbj|AB014075.1|[3868863]

Links

13: B88467 Reports

CpG0213B CpIOWAgDNA1 Cryptosporidium parvum genomic similar to TBP(TATA-
 binding protein)-interacting protein 49 (TIP49) homolog, genomic survey sequence
 gi|2970893|gb|B88467.1|[2970893]

Links

14: AI591696 Reports

vq30h01.y1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone
 IMAGE:1095793 5' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|4600744|gb|AI591696.1|[4600744]

Links

15: D84298 Reports

Dcinococcus radiodurans DNA for ruvB protein, complete cds
 gi|4586353|dbj|D84298.1|[4586353]

Links

16: AE000613 Reports

Helicobacter pylori 26695 section 91 of 134 of the complete genome
 gi|2314200|gb|AE000613.1|[AE000613][2314200]

Links

17: AI585221 Reports

fb95g04.y1 Zebrafish WashU MPI/IMG EST Danio rerio cDNA clone IMAGE:3719670 5'
 similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|4571118|gb|AI585221.1|[4571118]

Links

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- 18: [AI584574](#) Reports Links
fb95g04.x1 Zebrafish WashU MPIMG EST Danio rerio cDNA clone IMAGE:3719670 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4570471|gb|AI584574.1|[4570471]
- 19: [AI565821](#) Reports Links
tn20g08.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2168222 3' similar to SW:YEBA_ECOLI P24204 HYPOTHETICAL 46.7 KD PROTEIN IN MSBB-RUVB INTERGENIC REGION ;, mRNA sequence
gi|4524278|gb|AI565821.1|[4524278]
- 20: [AI539151](#) Reports Links
tp73c06.x1 NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2204938 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4453286|gb|AI539151.1|[4453286]
- 21: [AI503510](#) Reports Links
vq30h01.x1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone IMAGE:1095793 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4401361|gb|AI503510.1|[4401361]
- 22: [AI421332](#) Reports Links
tf14f12.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2096207 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4267263|gb|AI421332.1|[4267263]
- 23: [AI298410](#) Reports Links
qn01f05.x1 NCI_CGAP_Lu5 Homo sapiens cDNA clone IMAGE:1896993 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3958146|gb|AI298410.1|[3958146]
- 24: [AF064953](#) Reports Links
Coxiella burnetii Holliday junction DNA helicase RuvB gene, complete cds
gi|3248954|gb|AF064953.1|AF064953[3248954]
- 25: [AE001471](#) Reports Links
Helicobacter pylori, strain J99 section 32 of 132 of the complete genome
gi|4154880|gb|AE001471.1|AE001471[4154880]
- 26: [AI361999](#) Reports Links
qy49c08.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2015342 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4113620|gb|AI361999.1|[4113620]
- 27: [AI351057](#) Reports Links
qt22c06.x1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1948330 3' similar to

Entrez Nucleotide

TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|4088263|gb|AI351057.1|[4088263]

Links

 28: AF070735 Reports

Homo sapiens RuvB-like protein RUVBL1 mRNA, complete cds
gi|3243034|gb|AF070735.1|[3243034]

Links

 29: AI084169 Reports

qf27e09.x1 NCI_CGAP_Bm25 Homo sapiens cDNA clone IMAGE:1751272 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3422592|gb|AI084169.1|[3422592]

Links

 30: AI183800 Reports

qe18h08.x1 Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1739391 3'
similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3734438|gb|AI183800.1|[3734438]

Links

 31: AI096984 Reports

qb42d09.x1 NCI_CGAP_Bm23 Homo sapiens cDNA clone IMAGE:1702769 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3446566|gb|AI096984.1|[3446566]

Links

 32: AI078327 Reports

oz12d03.x1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone
IMAGE:1675109 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3412735|gb|AI078327.1|[3412735]

Links

 33: U55214 Reports

Treponema pallidum RuvB' gene, partial cds, and TroA, TroB, TroC, TroD, TroR, Pgm,
Pfs, and Tpp15 genes, complete cds
gi|1777931|gb|U55214.1|[TPU55214[1777931]]

Links

 34: AI033780 Reports

ox20a01.x1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone
IMAGE:1656840 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3254733|gb|AI033780.1|[3254733]

Links

 35: AA906726 Reports

ok78g03.s1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1520116 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3042312|gb|AA906726.1|[3042312]

Links

 36: AI049841 Reports

an21e02.x1 Gessler Wilms tumor Homo sapiens cDNA clone IMAGE:1699322 3' similar
to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3298958|gb|AI049841.1|[3298958]

Links

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- 37: [AA947109 Reports](#) Links
 om79a04.s1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:1553358 3' similar to
 TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|3110504|gb|AA947109.1|[3110504]
- 38: [AA974522 Reports](#) Links
 of13d01.s1 NCI_CGAP_Co12 Homo sapiens cDNA clone IMAGE:1420993 3' similar to
 TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|3149702|gb|AA974522.1|[3149702]
- 39: [AI006350 Reports](#) Links
 ua70f09.r1 Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1362857 5'
 similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|3215959|gb|AI006350.1|[3215959]
- 40: [AA985767 Reports](#) Links
 ua66e06.r1 Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1362466 5'
 similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|3167156|gb|AA985767.1|[3167156]
- 41: [AB012122 Reports](#) Links
 Homo sapiens mRNA for TIP49, complete cds
 gi|3132307|dbj|AB012122.1|[3132307]
- 42: [AA878287 Reports](#) Links
 oc57g05.s1 NCI_CGAP_Lu5 Homo sapiens cDNA clone IMAGE:1415768 3' similar to
 TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|2987252|gb|AA878287.1|[2987252]
- 43: [AA869108 Reports](#) Links
 vq30h01.r1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone
 IMAGE:1095793 5' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
 gi|2964553|gb|AA869108.1|[2964553]
- 44: [AA606258 Reports](#) Links
 vo47c01.r1 Barstead mouse irradiated colon MPLRB7 Mus musculus cDNA clone
 IMAGE:1053024 5' similar to SW:RUVB_HAEIN P44631 HOLLIDAY JUNCTION DNA
 HELICASE RUVB. ;, mRNA sequence
 gi|2454716|gb|AA606258.1|[2454716]
- 45: [D90829 Reports](#) Links
 E.coli genomic DNA, Kohara clone #337(41.9-42.3 min.)
 gi|1736504|dbj|D90829.1|[1736504]
- 46: [D90828 Reports](#) Links
 E.coli genomic DNA, Kohara clone #336gap(41.6-41.9 min.)
 gi|1736490|dbj|D90828.1|[1736490]

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- 47: [D83138](#) Reports Links
Pseudomonas aeruginosa ruvC, ruvA and ruvB genes for Holliday junction-related proteins
gi|1183837|dbj|D83138.1|PSERUVABC[1183837]
- 48: [U52110](#) Reports Links
Sulfolobus solfataricus putative ribokinase and Dbh genes, complete cds, and putative ATPase gene, partial cds
gi|1706950|gb|U52110.1|SSU52110[1706950]
- 49: [U38840](#) Reports Links
Thermotoga maritima RuvB protein (RUVB) gene, complete cds
gi|1063667|gb|U38840.1|TMU38840[1063667]
- 50: [X07091](#) Reports Links
E. coli ruv region
gi|42901|emb|X07091.1|ECRUVAB[42901]
- 51: [U38892](#) Reports Links
Synechocystis sp. ruvB gene, complete cds, and secA gene, partial cds
gi|1256587|gb|U38892.1|SSU38892[1256587]
- 52: [M86239](#) Reports Links
Cyanophora paradoxa cyanelle transfer RNA-Asn (trnN) gene, 3' end; transfer RNA-Leu (trnL) gene; FA/FB (psaC) gene, complete cds; ORF243, complete cds; ORF162, 3' end
gi|336634|gb|M86239.1|CPACYLNAA[336634]
- 53: [M21298](#) Reports Links
E.coli ruv operon (ruvA and ruvB) genes encoding proteins for filament formation and sensitivity to UV radiation, complete cds
gi|147781|gb|M21298.1|ECORUVABA[147781]
- 54: [U00011](#) Reports Links
Mycobacterium leprae cosmid B1177
gi|466807|gb|U00011.1|U00011[466807]

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